

WEST Search History

DATE: Thursday, July 21, 2005

Hide?	Set Name	Query	Hit Count
		<i>DB=PGPB,USPT,USOC,EPAB,JPAB,DWPI; PLUR=YES; OP=ADJ</i>	
<input type="checkbox"/>	L1	tamura-T\$.in.	9140
<input type="checkbox"/>	L2	oto-N\$.in. or suzuki-N\$.in. or mizuno-K\$.in. L1	13433
<input type="checkbox"/>	L3	oto-N\$.in. or suzuki-N\$.in. or mizuno-K\$.in.	17929
<input type="checkbox"/>	L4	yamamoto-N\$.in. or suzuki-N\$.in. or mizuno-K\$.in.	25144
<input type="checkbox"/>	L5	(L1 or L4) and (biochip or array)	704
<input type="checkbox"/>	L6	(L1 or L4) and (biochip or microarray or array)	723
<input type="checkbox"/>	L7	L6 and hybridiz\$	66
<input type="checkbox"/>	L8	L7 and biopolymer	21
<input type="checkbox"/>	L9	L8 and visual\$	7
<input type="checkbox"/>	L10	probe similarity score	0
<input type="checkbox"/>	L11	similarity score	1284
<input type="checkbox"/>	L12	biochip or microarray	18908
<input type="checkbox"/>	L13	L12 same bipolymer	1
<input type="checkbox"/>	L14	L11 and L12	77
<input type="checkbox"/>	L15	L14 and biopolymer	16
<input type="checkbox"/>	L16	L15 and (imag\$ and visual\$)	11
<input type="checkbox"/>	L17	L16 and hybridiz\$	11
<input type="checkbox"/>	L18	L12 and biopolymer	1868
<input type="checkbox"/>	L19	L18 and hybridiz\$	1657
<input type="checkbox"/>	L20	L19 and (similar\$ scor\$ or percent homology or percent identity or sequence identity)	555
<input type="checkbox"/>	L21	L20 and (display\$ and imag\$ and visual\$)	333
<input type="checkbox"/>	L22	L21 and (different same color)	30
<input type="checkbox"/>	L23	L22 and (different same (valu\$ or depth))	23
<input type="checkbox"/>	L24	L23 and spot imag\$	1
<input type="checkbox"/>	L25	L23 and statistic\$	23
<input type="checkbox"/>	L26	L25 and matrix	21
<input type="checkbox"/>	L27	L26 and (hybridization level or hybridization profile)	1
<input type="checkbox"/>	L28	L25 and storage unit	1
<input type="checkbox"/>	L29	L25 and (data near storage)	7

<input type="checkbox"/>	L30	L25 and hybridization	23
<input type="checkbox"/>	L31	((biopolymer or probe) near (biochip or microarray))	227
<input type="checkbox"/>	L32	L31 and ((display\$ or visual\$) same hybridiz\$)	17
<input type="checkbox"/>	L33	(display\$ or visual\$) same (percent identity or percent homology or sequence identity or sequence similarity or similarity score)	3442
<input type="checkbox"/>	L34	L33 same (hybridization)	116
<input type="checkbox"/>	L35	L34 and (graphic\$)	14
<input type="checkbox"/>	L36	L33 and (different near color)	64
<input type="checkbox"/>	L37	L36 and (image)	39
<input type="checkbox"/>	L38	L31 and L33	25
<input type="checkbox"/>	L39	(display\$ or visual\$)near(percent identity or percent homology or sequence identity or sequence similarity or similarity score)	144
<input type="checkbox"/>	L40	L39 and hybridiz\$	115
<input type="checkbox"/>	L41	L40 and (biochip or microarray)	26
<input type="checkbox"/>	L42	L41 and (biopolymer or probe)	26
<input type="checkbox"/>	L43	L39 and (biochip or microarray)	26
<input type="checkbox"/>	L44	L33 and (biochip or microarray)	533
<input type="checkbox"/>	L45	L44 and hybridization	497
<input type="checkbox"/>	L46	L45 and ((Imag\$ or matrix or graphic\$) same (different near color))	1
<input type="checkbox"/>	L47	L45 and ((Imag\$ or matrix or graphic\$) and (different near color))	6
<input type="checkbox"/>	L48	BLAST same (imag\$ or graphic\$ or visual\$)	4587
<input type="checkbox"/>	L49	L48 same (biochip or microarray)	5
<input type="checkbox"/>	L50	6188783.pn. or 5812272.pn. or 6471916.pn. or 6284465.pn.	8
<input type="checkbox"/>	L51	6349144.pn. or 4852183.pn. or 6306643.pn. or 6228575.pn.	8
<input type="checkbox"/>	L52	(identity or similarity) near score	2501
<input type="checkbox"/>	L53	L52 same probe\$	12
<input type="checkbox"/>	L54	L52 and L48	29
<input type="checkbox"/>	L55	L54 and prob\$	28
<input type="checkbox"/>	L56	L52 and (BLAST or smith-watson or algorithm)	2036
<input type="checkbox"/>	L57	L52 same (probe or oligonucleotide or biopolymer or oligomer)	19
<input type="checkbox"/>	L58	L57 and @pd > 20050202	2

END OF SEARCH HISTORY